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(54) **ORDERED BIOLOGICAL
NANOSTRUCTURES FORMED FROM
CHAPERONIN POLYPEPTIDES**

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C07K 14/00 (2006.01)
A61K 38/00 (2006.01)

(52) **U.S. Cl.** **530/350; 514/2; 977/773**

(58) **Field of Classification Search** None
See application file for complete search history.

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(57) **ABSTRACT**

The following application relates to nanotemplates, nano-
structures, nanoarrays and nanodevices formed from wild-
type and mutated chaperonin polypeptides, methods of pro-
ducing such compositions, methods of using such
compositions and particular chaperonin polypeptides that can
be utilized in producing such compositions.

17 Claims, 36 Drawing Sheets

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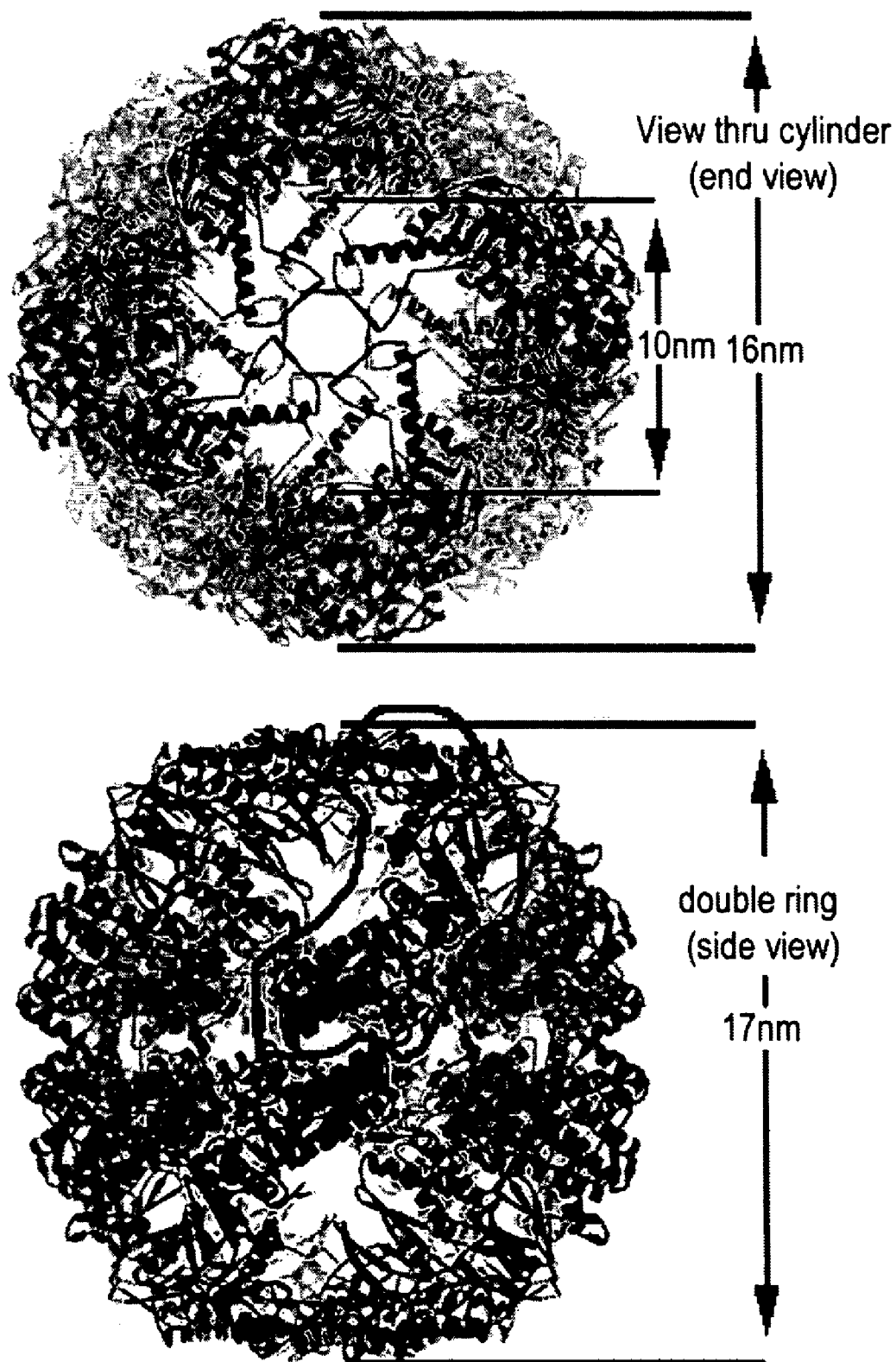


FIG.1 Prior Art

[illegible]

TF55 beta - *S. shibatae*
GroEL - *E. coli*
thermosome beta - *T. acidophilum*
cyanobacterial HSP60 *synechococcus*
HSP60-4 *M. acetivorans*
HSP65 - *M. tuberculosis*
thermosome alpha - *A. pernix*
thermosome alpha - *M. mazei*
mitochondrial HSP60 - *A. thaliana*
TCP1 alpha - YEAST
mitochondrial HSP60 - HUMAN
mitochondrial HSP60 - MOUSE
TCP1 alpha - HUMAN
TCP1 alpha - MOUSE
Consensus

LEGEND:

- ☒ IDENTICAL RESIDUES EXA.
- ☒ BLOCK OF SIMILAR EXA.
- ☒ CONSERVATIVE EXA.
- ☒ WEAKLY SIMILAR EXA.
- ☐ NON-SIMILAR EXA.

FIG. 2A

		Section 2										68	

		Section 3																																							
		102																																							
		90																																							
		80																																							
		69																																							
TF55 beta - <i>S. shibatae</i>		(57)	M	F	V	D	S	L	G	D	I	T	I	T	N	D	G	A	T	I	L	D	K	M	D	L	Q	H	P	-	-	-	-	T	G	K					
GroEL - <i>E. coli</i>		(39)	V	L	D	K	S	F	G	A	P	T	I	T	K	D	G	V	S	V	A	R	E	I	E	L	E	D	K	F	E	N	M	G	A	Q					
thermosome beta - <i>T. acidophilum</i>		(49)	M	L	V	D	S	L	G	D	I	V	I	T	N	D	G	V	T	I	L	K	E	M	D	V	E	H	P	-	-	-	-	A	A	K					
cyanobacterial HSP60 synecchococcus		(38)	V	L	E	K	K	F	G	A	P	Q	I	T	N	D	G	V	T	I	A	K	E	I	E	L	E	D	H	I	E	N	T	G	V	A					
HSP60-4 <i>M. acetivorans</i>		(68)	I	I	V	N	P	V	G	D	I	F	V	T	S	D	G	K	V	I	L	K	E	I	D	V	L	H	P	-	-	-	-	I	V	T					
HSP65 - <i>M. tuberculosis</i>		(38)	V	L	E	K	K	W	G	A	P	T	I	T	N	D	G	V	S	I	A	K	E	I	E	L	E	D	P	Y	E	K	I	G	A	E					
thermosome alpha - <i>A. pernix</i>		(50)	M	L	V	D	A	F	G	D	I	T	V	T	N	D	G	A	T	I	V	K	E	M	E	I	Q	H	P	-	-	-	-	A	A	K					
thermosome alpha - <i>M. mazei</i>		(48)	M	L	V	D	A	M	G	D	V	V	I	T	N	D	G	A	T	I	L	K	E	M	D	I	E	H	P	-	-	-	-	C	A	K					
mitochondrial HSP60 - <i>A. thaliana</i>		(69)	V	I	E	Q	S	W	G	A	P	K	V	T	K	D	G	V	V	A	K	S	I	E	F	K	D	K	I	K	N	V	G	A	S						
TCP1 alpha - YEAST		(52)	M	L	V	D	D	I	G	D	F	T	V	T	N	D	G	A	T	I	L	S	L	D	V	L	Q	H	P	-	-	-	-	A	G	K					
mitochondrial HSP60 - HUMAN		(52)	-	A	K	R	S	Y	G	-	-	-	-	-	-	-	-	-	Q	L	P	P	S	L	A	L	Q	D	K	Y	K	N	T	G	A	K					
mitochondrial HSP60 - MOUSE		(63)	I	I	E	Q	S	W	G	S	P	K	V	T	K	D	G	V	T	V	A	K	S	I	D	L	K	D	K	Y	K	N	I	G	A	K					
TCP1 alpha - HUMAN		(44)	M	L	V	D	D	I	G	D	V	T	I	T	N	D	G	A	T	I	L	K	L	L	E	V	E	H	P	-	-	-	-	A	A	K					
TCP1 alpha - MOUSE		(44)	M	L	V	D	D	I	G	D	V	T	I	T	N	D	G	A	T	I	L	K	L	L	E	V	E	H	P	-	-	-	-	A	A	K					
Consensus		(69)	M	L	V	D	S	W	G	D	I	T	I	T	N	D	G	-	-	-	-	-	-	T	I	L	K	E	I	E	L	E	H	P	-	-	-	-	G	A	K

FIG.2C

LEGEND:

- G IDENTICAL RESIDUES EXA.
- G BLOCK OF SIMILAR EXA.
- G CONSERVATIVE EXA.
- G** WEAKLY SIMILAR EXA.
- G NON-SIMILAR EXA.

Section 4

	(103)	103	110	120	136
TF55 beta - <i>S. shibatae</i>	(87)	L	V Q I A K G Q D E	T A D G T K T A V I L A G E L A K A E D L	
GroEL - <i>E. coli</i>	(73)	M V	K E V A S K A N D A A	C D G T T T A T V L A Q A I I T E G L K A	
thermosome beta - <i>T. acidophilum</i>	(79)	M M	V E V S K T Q D S F V	G D G T T T A V I I A G L Q Q A Q G L	
cyanobacterial HSP60 synecchococcus	(72)	L I I R Q A A S K T N D A A	G D G T T T A T V L A H A V V K E G L R N		
HSP60-4 <i>M. acetivorans</i>	(98)	S L K K L A E S M D K A C G D G T K T A V I F A S N L I I K N A V R L			
HSP65 - <i>M. tuberculosis</i>	(72)	L V K E V A K K T I D V A	G D G T T T A T V L A Q A L V R E G L R N		
thermosome alpha - <i>A. pernix</i>	(80)	L L V E V A K A Q D A E	G D G T T T A V V L A G A L L E K A E K L		
thermosome alpha - <i>M. mazei</i>	(78)	M I V E V A K T Q D A E	G D G T T T A V V L A G A L L T K A E D L		
mitochondrial HSP60 - <i>A. thaliana</i>	(103)	L V K Q V A N A T N D V A	G D G T T T C A T V L T R A I F A E G C K S		
TCP1 alpha - YEAST	(82)	I L V E L A Q Q D R E I	G D G T T T S V I I I A S E L L K R A N E L		
mitochondrial HSP60 - HUMAN	(76)	L V Q D V A N N T N E E A V	D G T T T V T A L A R S I A K E G F E K		
mitochondrial HSP60 - MOUSE	(97)	L V Q D V A N N T N E E A G	D G T T T S T V L A R S I A K E G F E K		
TCP1 alpha - HUMAN	(74)	V L C E L A D L Q D K E	G D G T T T S V I I I A A E L L K N A D E L		
TCP1 alpha - MOUSE	(74)	V L C E L A D L Q D K E	G D G T T T S V I I I A A E L L K N A D E L		
Consensus	(103)	L L E V A Q D D E G D G T T T A V V L A A L L K A E L			

LEGEND:

- G IDENTICAL RESIDUES EXA.
- G BLOCK OF SIMILAR EXA.
- G CONSERVATIVE EXA.
- G WEAKLY SIMILAR EXA.
- G NON-SIMILAR EXA.

FIG.2D

		Section 5									
		170									
		160									
		150									
		140									
		130									
		120									
		110									
		100									
		90									
		80									
		70									
		60									
		50									
		40									
		30									
		20									
		10									
		0									

	Section 6																																			
	(171) 171							180							190							204														
TF55 beta - <i>S. shibatae</i>	(154)	I	N	D	T	D	V	L	R	K	V	A	L	T	S	L	G	S	K	A	V	A	G	-	-	A	R	E	Y	L	A	D	L	V	V	
GroEL - <i>E. coli</i>	(138)	C	S	D	S	K	A	I	A	Q	V	G	T	I	S	A	N	S	D	E	T	V	G	K	L	I	A	E	A	M	D	K	V	G	K	
thermosome beta - <i>Tacidophilum</i>	(146)	A	D	E	K	A	L	L	K	M	A	Q	T	S	L	N	S	K	S	A	S	V	-	-	A	K	D	K	L	A	E	I	S	Y		
cyanobacterial HSP60 synecococcus	(137)	V	E	D	S	K	S	I	A	Q	V	G	A	I	S	A	T	G	N	D	F	E	V	G	Q	M	I	A	D	A	M	D	K	V	G	K
HSP60-4 <i>M. acetivorans</i>	(163)	-	A	S	E	E	D	I	R	T	T	I	M	C	S	A	T	G	K	I	E	R	Q	-	Q	A	Q	A	A	V	T	E	I	A	L	
HSP65 - <i>M. tuberculosis</i>	(137)	V	E	T	K	E	Q	I	A	A	T	A	I	S	A	I	G	-	D	Q	S	I	G	D	L	I	A	E	A	M	D	K	V	G	N	
thermosome alpha - <i>A. pernix</i>	(147)	V	E	D	S	V	L	R	R	T	A	E	T	I	A	G	T	L	A	S	K	F	V	G	T	G	P	E	R	D	I	S	M	V	I	
thermosome alpha - <i>M. maezi</i>	(145)	P	E	D	T	E	T	L	E	K	I	A	G	T	I	A	I	T	G	K	G	A	E	S	-	-	H	K	A	H	L	S	N	L	A	V
mitochondrial HSP60 - <i>A. thaliana</i>	(168)	I	S	T	S	E	E	I	A	Q	V	G	T	I	S	A	N	G	E	R	E	I	G	E	L	I	A	K	A	M	E	K	V	G	K	
TCP1 alpha - YEAST	(150)	T	L	G	K	E	T	L	I	N	I	A	I	K	T	S	M	S	S	K	I	I	G	A	D	-	-	S	D	F	S	N	M	V	V	
mitochondrial HSP60 - HUMAN	(141)	V	T	T	P	E	E	I	A	R	V	A	T	I	S	A	N	G	D	K	E	I	G	N	I	I	S	D	A	M	K	K	V	G	S	
mitochondrial HSP60 - MOUSE	(162)	V	T	T	P	E	E	I	A	Q	V	A	T	I	S	A	N	G	D	K	D	I	G	N	I	I	S	D	A	M	K	K	V	G	R	
TCP1 alpha - HUMAN	(142)	E	L	G	R	D	C	L	I	N	A	A	K	T	S	M	S	S	K	I	I	G	I	N	-	-	G	D	F	A	N	M	V	V	V	
TCP1 alpha - MOUSE	(142)	E	L	G	R	D	C	L	I	N	A	A	K	T	S	M	S	S	K	I	I	G	I	N	-	-	G	D	F	A	N	M	V	V	V	
Consensus	(171)	V				E	I	Q	V	A	T	S	A	S	K									G			A	D	A	M				V	G	V

LEGEND:

[G] IDENTICAL RESIDUES EXA.

[G] BLOCK OF SIMILAR EXA.

[G] CONSERVATIVE EXA.

G WEAKLY SIMILAR EXA.

G NON-SIMILAR EXA.

FIG. 2F

Section 8

	(239)	239	250	260	272
TF55 beta - <i>S. shibatae</i>	(219)	T Q L V Y G I V V D K E V V H P P G M P K R I E N - - A K I A L D A			
GroEL - <i>E. coli</i>	(206)	N K P E T G A V E L E S P F I L L A D K K I S N I R E M L P V L E A			
thermosome beta - <i>T. acidophilum</i>	(211)	T Q L I N G I I V D K E K V H P P G M P D V V I K D - - A K I A L D A			
cyanobacterial HSP60	(205)	I D T E R M E A V F D E N P A R E D M P K I S Y Q N P A V L I T N Y D L			
synecchococcus	(225)	I V A I E G L I M D E N P A R E D M P K I S Y Q N P A V L I T N Y D L			
HSP60-4 <i>M. acetivorans</i>	(204)	T D P E R Q E A V L E D P Y I L L V S S K V S T V K D L L P L I E K			
HSP65 - <i>M. tuberculosis</i>	(215)	S K L V R G I V L D K E V V H P P A M P K R V E N - - A K I L V L D A			
thermosome alpha - <i>A. pernix</i>	(211)	S E I V E G V I V D K E R V H T G M P E V V I K D - - A K I V L L S V			
thermosome alpha - <i>M. mazei</i>	(236)	T N Q K T Q K C E L D D P L I L I H E K K I S S I N S I V K V L E L			
mitochondrial HSP60 - <i>A. thaliana</i>	(216)	S L V P G Y A L N C T V A S Q A M P K R I I A G G N V K I A C L D L			
TCP1 alpha - YEAST	(185)	- - - S Q K C E F Q D A Y V L L S E K K I I S S V Q Q S I A P A I E I			
mitochondrial HSP60 - HUMAN	(230)	N T S K G Q K C E F Q D A Y V L L S E K K I I S S V Q Q S I A P A I E I			
mitochondrial HSP60 - MOUSE	(208)	S M L I S G Y A L N C V V G S Q G M P K R I I V N - - A K I A C L D F			
TCP1 alpha - HUMAN	(208)	S M L I S G Y A L N C V V G S Q G M P K R I I V N - - A K I A C L D F			
TCP1 alpha - MOUSE	(208)	S M L I S G Y A L N C V V G S Q G M P K R I I V N - - A K I A C L D F			
Consensus	(239)	S L I G V E G M P K K I A K I L L D			

LEGEND:

[G] IDENTICAL RESIDUES EXA.

[G] BLOCK OF SIMILAR EXA.

[G] CONSERVATIVE EXA.

G WEAKLY SIMILAR EXA.

G NON-SIMILAR EXA.

FIG. 2H